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## RAW SEQUENCE LISTING

DATE: 09/18/2001

PATENT APPLICATION: US/09/938,641

TIME: 11:13:51

Input Set : A:\211226US0X.txt

Output Set: N:\CRF3\09182001\I938641.raw

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3 <110> APPLICANT: MARX, ACHIM
4     FARWICK, MIKE
5     HERMANN, THOMAS
6     SCHISCHKA, NATALIE
7     BATHE, BRIGITTE
9 <120> TITLE OF INVENTION: NUCLEOTIDE SEQUENCES WHICH CODE FOR THE oxyR GENE
11 <130> FILE REFERENCE: 211226US0X
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/938,641
C--> 13 <141> CURRENT FILING DATE: 2001-08-27
13 <150> PRIOR APPLICATION NUMBER: DE 10042052.4
14 <151> PRIOR FILING DATE: 2000-08-26
16 <150> PRIOR APPLICATION NUMBER: DE 10110053.1
17 <151> PRIOR FILING DATE: 2001-03-02
19 <150> PRIOR APPLICATION NUMBER: US 60/279,415
20 <151> PRIOR FILING DATE: 2001-03-29
22 <160> NUMBER OF SEQ ID NOS: 4
24 <170> SOFTWARE: PatentIn version 3.1
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27 <211> LENGTH: 1675
28 <212> TYPE: DNA
29 <213> ORGANISM: Corynebacterium glutamicum
31 <220> FEATURE:
32 <221> NAME/KEY: CDS
33 <222> LOCATION: (491)..(1471)
34 <223> OTHER INFORMATION:
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38 gccaacgcga gggcatttac catcatggtg cgcaacgccg tgttccgcct tgtggagcta      60
40 tttgcttatg aaaaggaaga tcagcttagt cagatgactg aatacctgga tgaggctcct      120
42 gatttcggtg ctgcgatgga tgcgtacttt gatgaatatg cggatcttga taccggccccg      180
44 gcagctcgtg gaaccagagtt cttcaaggta gagcacacgg gaagaatgtg ggagggtgcgt      240
46 cagggtgtga aggatccaga aggtgataat tccttcgcgt ttgttgccac cattgatctt      300
48 gatgcctctg atgatgcagg tgagggtgcgt tttggatcgc tgcgattga ccacaactag      360
50 gggtttgctg cgaaaagcaa gcacgcctgg tgctgattt gagcggtttt acctatggcg      420
52 ctttggcgcc gtcaaaactgt cccagcgatt tcattattat tttcgtgcat tcaccgttat      480
54 agttataggg atg agc aat aaa gag tac cgg ccc aca ctc gcc cag ctt      529
55     Met Ser Asn Lys Glu Tyr Arg Pro Thr Leu Ala Gln Leu
56     1             5             10
58 cgc acc ttt gtc acc atc gca gaa tgc aag cac ttt ggt act gct gcc      577
59 Arg Thr Phe Val Thr Ile Ala Glu Cys Lys His Phe Gly Thr Ala Ala
60     15             20             25
62 acc aag ctg tcc att tcg cag cca tcc ctc tcc cag gca ctt gtc gca      625
63 Thr Lys Leu Ser Ile Ser Gln Pro Ser Leu Ser Gln Ala Leu Val Ala
64 30             35             40             45
66 tta gaa aca ggc ctg gga gtt cag ctg att gaa cgc tcc acc cgc aag      673
67 Leu Glu Thr Gly Leu Gly Val Gln Leu Ile Glu Arg Ser Thr Arg Lys
68     50             55             60
70 gtc att gtc acc cca gcg ggc gag aag ttg ctg cca ttc gcc aaa tcc      721

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71 Val Ile Val Thr Pro Ala Gly Glu Lys Leu Leu Pro Phe Ala Lys Ser
72          65          70          75
74 acc ctt gac gcg gcg gag tct ttc ctc tcc cac gcc aag ggc gcc aac      769
75 Thr Leu Asp Ala Ala Glu Ser Phe Leu Ser His Ala Lys Gly Ala Asn
76          80          85          90
78 ggt tcg ctc act gga ccg ttg acc gta ggc atc atc ccc acg gcg gct      817
79 Gly Ser Leu Thr Gly Pro Leu Thr Val Gly Ile Ile Pro Thr Ala Ala
80          95          100          105
82 cct tac att ttg ccg tca atg ctg tcc atc gtg gat gaa gaa tat cca      865
83 Pro Tyr Ile Leu Pro Ser Met Leu Ser Ile Val Asp Glu Glu Tyr Pro
84 110          115          120          125
86 gat ctg gaa cct cac atc gtc gag gac caa acc aag cat ctt ctc gcg      913
87 Asp Leu Glu Pro His Ile Val Glu Asp Gln Thr Lys His Leu Leu Ala
88          130          135          140
90 ttg ctg cgc gac ggc gcc atc gac gtc gcc atg atg gcc ctg cct tct      961
91 Leu Leu Arg Asp Gly Ala Ile Asp Val Ala Met Met Ala Leu Pro Ser
92          145          150          155
94 gag gca cca ggc atg aag gaa atc ccc ctc tac gac gaa gac ttt atc      1009
95 Glu Ala Pro Gly Met Lys Glu Ile Pro Leu Tyr Asp Glu Asp Phe Ile
96          160          165          170
98 gtc gtt aca gct agc gat cac ccc ttc gcc ggc cgc caa gac tta gaa      1057
99 Val Val Thr Ala Ser Asp His Pro Phe Ala Gly Arg Gln Asp Leu Glu
100          175          180          185
102 cta tcc gcc tta gaa gac ctc gat ctg ctg ctt ctc gac gac gga cac      1105
103 Leu Ser Ala Leu Glu Asp Leu Asp Leu Leu Leu Leu Asp Asp Gly His
104 190          195          200          205
106 tgc ctc cac gac caa att gtg gac ctg tgc cgc cgc gga gac atc aac      1153
107 Cys Leu His Asp Gln Ile Val Asp Leu Cys Arg Arg Gly Asp Ile Asn
108          210          215          220
110 ccc att agc tcc act act gct gtc acc cgc gca tcc agc ctt acc acc      1201
111 Pro Ile Ser Ser Thr Thr Ala Val Thr Arg Ala Ser Ser Leu Thr Thr
112          225          230          235
114 gtc atg cag ctc gtc gtc gcc ggc ctt gga tcc acc ttg gtc cca atc      1249
115 Val Met Gln Leu Val Val Ala Gly Leu Gly Ser Thr Leu Val Pro Ile
116          240          245          250
118 agc gca atc cca tgg gaa tgc acc cga cca gga ctg gca aca gcc aac      1297
119 Ser Ala Ile Pro Trp Glu Cys Thr Arg Pro Gly Leu Ala Thr Ala Asn
120          255          260          265
122 ttc aac tct gat gtc acc gca aac cgc cgc att gga ttg gtg tac cgt      1345
123 Phe Asn Ser Asp Val Thr Ala Asn Arg Arg Ile Gly Leu Val Tyr Arg
124 270          275          280          285
126 tcc tct tct tct cgc gcc gaa gag ttc gaa cag ttt gca ctc att ttg      1393
127 Ser Ser Ser Ser Arg Ala Glu Glu Phe Glu Gln Phe Ala Leu Ile Leu
128          290          295          300
130 cag cgc gct ttc caa gaa gcc gtc gcg ctt gct gcc tca act ggc atc      1441
131 Gln Arg Ala Phe Gln Glu Ala Val Ala Leu Ala Ala Ser Thr Gly Ile
132          305          310          315
134 acc ttg aag caa aat gtc gcg gta gcg cag taagtttttc tagaggtttt      1491
135 Thr Leu Lys Gln Asn Val Ala Val Ala Gln

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136          320          325
138 ccagagtcag ctacaagcaa aaagcccttt ccattgatgc acaccaacgt gagattcaag 1551
140 ggaaagggtc ttattgattg cagaatgcct actgcattag cggcgctcca ccggaatatt 1611
142 tccaccactg atctggcggt aaatatgaac ggtagacagc atcattactg gcagcacgat 1671
144 gatc 1675
147 <210> SEQ ID NO: 2
148 <211> LENGTH: 327
149 <212> TYPE: PRT
150 <213> ORGANISM: Corynebacterium glutamicum
152 <400> SEQUENCE: 2
154 Met Ser Asn Lys Glu Tyr Arg Pro Thr Leu Ala Gln Leu Arg Thr Phe
155 1 5 10 15
158 Val Thr Ile Ala Glu Cys Lys His Phe Gly Thr Ala Ala Thr Lys Leu
159 20 25 30
162 Ser Ile Ser Gln Pro Ser Leu Ser Gln Ala Leu Val Ala Leu Glu Thr
163 35 40 45
166 Gly Leu Gly Val Gln Leu Ile Glu Arg Ser Thr Arg Lys Val Ile Val
167 50 55 60
170 Thr Pro Ala Gly Glu Lys Leu Leu Pro Phe Ala Lys Ser Thr Leu Asp
171 65 70 75 80
174 Ala Ala Glu Ser Phe Leu Ser His Ala Lys Gly Ala Asn Gly Ser Leu
175 85 90 95
178 Thr Gly Pro Leu Thr Val Gly Ile Ile Pro Thr Ala Ala Pro Tyr Ile
179 100 105 110
182 Leu Pro Ser Met Leu Ser Ile Val Asp Glu Glu Tyr Pro Asp Leu Glu
183 115 120 125
186 Pro His Ile Val Glu Asp Gln Thr Lys His Leu Leu Ala Leu Leu Arg
187 130 135 140
190 Asp Gly Ala Ile Asp Val Ala Met Met Ala Leu Pro Ser Glu Ala Pro
191 145 150 155 160
194 Gly Met Lys Glu Ile Pro Leu Tyr Asp Glu Asp Phe Ile Val Val Thr
195 165 170 175
198 Ala Ser Asp His Pro Phe Ala Gly Arg Gln Asp Leu Glu Leu Ser Ala
199 180 185 190
202 Leu Glu Asp Leu Asp Leu Leu Leu Asp Asp Gly His Cys Leu His
203 195 200 205
206 Asp Gln Ile Val Asp Leu Cys Arg Arg Gly Asp Ile Asn Pro Ile Ser
207 210 215 220
210 Ser Thr Thr Ala Val Thr Arg Ala Ser Ser Leu Thr Thr Val Met Gln
211 225 230 235 240
214 Leu Val Val Ala Gly Leu Gly Ser Thr Leu Val Pro Ile Ser Ala Ile
215 245 250 255
218 Pro Trp Glu Cys Thr Arg Pro Gly Leu Ala Thr Ala Asn Phe Asn Ser
219 260 265 270
222 Asp Val Thr Ala Asn Arg Arg Ile Gly Leu Val Tyr Arg Ser Ser Ser
223 275 280 285
226 Ser Arg Ala Glu Glu Phe Glu Gln Phe Ala Leu Ile Leu Gln Arg Ala
227 290 295 300
230 Phe Gln Glu Ala Val Ala Leu Ala Ala Ser Thr Gly Ile Thr Leu Lys

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234 Gln Asn Val Ala Val Ala Gln
235          325
238 <210> SEQ ID NO: 3
239 <211> LENGTH: 30
240 <212> TYPE: DNA
241 <213> ORGANISM: Artificial Sequence
243 <220> FEATURE:
244 <223> OTHER INFORMATION: Synthetic DNA
246 <400> SEQUENCE: 3
247 gatcgagaat tcaaaggaag atcagcttag          30
250 <210> SEQ ID NO: 4
251 <211> LENGTH: 20
252 <212> TYPE: DNA
253 <213> ORGANISM: Artificial Sequence
255 <220> FEATURE:
256 <223> OTHER INFORMATION: Synthetic DNA
258 <400> SEQUENCE: 4
259 ggaaaacctc tagaaaaact          20
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/938,641

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Input Set : A:\211226US0X.txt

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L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date